SEQUENCE SUBMISSION

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10	SEQ ID NO: 7 provides primate IL-1RA polypeptide sequence. SEQ ID NO: 8 provides rodent IL-1\(\gamma\) (IGIF) polypeptide sequence. SEQ ID NO: 9 provides primate IL-1\(\gamma\) (IGIF) polypeptide sequence. SEQ ID NO: 10 provides rodent IL-1\(\epsilon\) polypeptide sequence. SEQ ID NO: 11 provides rodent IL-1\(\epsilon\) (Dypeptide sequence.	
15	SEQ ID NO: 12 provides primate IL-1R6 nucleotide sequence. SEQ ID NO: 13 provides primate IL-1R6 polypeptide sequence. SEQ ID NO: 14 provides rodent IL-1R6 nucleotide sequence. SEQ ID NO: 15 provides rodent IL-1R6 polypeptide sequence.	
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- Asn Pro Glu Met Cys Leu Tyr Cys Glu Lys Val Gly Glu Gln Pro Thr 10 $\,$ 85 $\,$ 90 $\,$ 95
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5 Pro Asn Val Asn Leu Glu Glu Lys Ile Asp Val Val Pro Ile Glu Pro
35 40 45

His Ala Leu Phe Leu Gly Ile His Gly Gly Lys Met Cys Leu Ser Cys 50 60

Val Lys Ser Gly Asp Glu Thr Arg Leu Gln Leu Glu Ala Val Asn Ile 65 70 75 80

Thr Asp Leu Ser Glu Asn Arg Lys Gln Asp Lys Arg Phe Ala Phe Ile 85 90 95

Arg Ser Asp Ser Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro $100 \\ 105 \\ 110$

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Tyr Met Tyr Lys Asp Ser Glu Val Arg Gly Leu Ala Val Thr Leu Ser $50 \\ 50 \\ 50$

Val Lys Asp Ser Lys Met Ser Thr Leu Ser Cys Lys Asn Lys Ile Ile 65 70 75 80

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Asp Leu Ile Phe Phe Gln Lys Arg Val Pro Gly His Asn Lys Met Glu $100 \hspace{1cm} 105 \hspace{1cm} 110 \hspace{1cm} 110 \hspace{1cm}$

Phe Glu Ser Ser Leu Tyr Glu Gly His Phe Leu Ala Cys Gln Lys Glu

115 120 125

Asp Asp Ala Phe Lys Leu Ile Leu Lys Lys Lys Asp Glu Asn Gly Asp 130 135 140

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Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile $50 \hspace{1cm} 55 \hspace{1cm} 60 \hspace{1cm}$

Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile 65 70 75 80

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Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys 100 105 110

Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu 115 120 125

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 Mus sp.

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Ser Ala Phe His Ser Thr Glu Thr Ile Val Asp Gly Lys Leu Tyr Asp 5 Ala Tyr Val Leu Tyr Pro Lys Pro His Lys Glu Ser Gln Arg His Ala Val Asp Ala Leu Val Leu Asn Ile Leu Pro Glu Val Leu Glu Arg Gln 10 Cys Gly Tyr Lys Leu Phe Ile Phe Gly Arg Asp Glu Phe Pro Gly Gln 425 Ala Val Ala Asn Val Ile Asp Glu Asn Val Lys Leu Cys Arg Arg Leu 15 440 Ile Val Ile Val Val Pro Glu Ser Leu Gly Phe Gly Leu Leu Lys Asn 450 455 460 20 Leu Ser Glu Glu Gln Ile Ala Val Tyr Ser Ala Leu Ile Gln Asp Gly Met Lys Val Ile Leu Ile Glu Leu Glu Lys Ile Glu Asp Tyr Thr Val 490 25 Met Pro Glu Ser Ile Gln Tyr Ile Lys Gln Lys His Gly Ala Ile Arg 505 Trp His Gly Asp Phe Thr Glu Gln Ser Gln Cys Met Lys Thr Lys Phe 30 Trp Lys Thr Val Arg Tyr His Met Pro Pro Arg Arg Cys Arg Pro Phe 535 35 Leu Arg Ser Thr Cys Arg Ser Thr His Leu Cys Thr Ala Pro Gln Ala 545 550 555 Gln Asn 40 <210> 14 <211> 1686 45 <212> DNA <213> Unknown Organism <220> <223> Description of Unknown Organism: rodent; surmised 50 Rattus sp. <220> <221> CDS <222> (1)..(1683) 55 <400> 14 atg ggg atg cca ccc ttg ctc ttc tgt tgg gtg tct ttc gtg ctt cca Met Gly Met Pro Pro Leu Leu Phe Cys Trp Val Ser Phe Val Leu Pro 5 10 15 60

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	Leu	Phe	Val	Ala 20	Ala	Gly	Asn	Cys	Thr 25	Asp	Val	Tyr	Met	His 30	His	Glu		
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1.5	agc Ser 65	cca Pro	atc Ile	tcc Ser	atc Ile	aac Asn 70	aga Arg	cac His	gtt Val	aga Arg	att Ile 75	cac His	cag Gln	gac Asp	cag Gln	tcc Ser 80	240	
15													ggc Gly				288	
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25													gaa Glu 125				384	
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35	ggc Gly 145	agt Ser	ctg Leu	acg Thr	tgc Cys	cat His 150	ctc Leu	tac Tyr	ttc Phe	cca Pro	gag Glu 155	agc Ser	tgt Cys	gtt Val	ttg Leu	gat Asp 160	480	
33													gtg Val				528	
40													atc Ile				576	
45													cac His 205				624	
50													aag Lys				672	
55													aac Asn				720	
JO													aat Asn				768	
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5						aac Asn											864
10	acc Thr	aat Asn 290	ctg Leu	tct Ser	ctg Leu	agg Arg	aat Asn 295	cac His	att Ile	ctg Leu	tac Tyr	aca Thr 300	gtg Val	aac Asn	ata Ile	aca Thr	912
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15						gcc Ala											1008
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30						ttc Phe											1152
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45						gcc Ala											1344
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10	Glu	Val	Gln	Leu	Gly 245	Ser	Thr	Leu	Ile	Val 250	Asp	Cys	Asn	Ile	Thr 255	Ası
15	Thr	Lys	Glu	Asn 260	Thr	Asn	Leu	Arg	Cys 265	Trp	Arg	Val	Àsn	Asn 270	Thr	Let
	Val	Asp	Asp 275	Tyr	Tyr	Asn	Asp	Phe 280	Lys	Arg	Ile	Gln	Glu 285	Gly	Ile	Glu
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	Phe 305	Leu	Glu	Va1	Lys	Met 310	Glu	Asp	Tyr	Gly	His 315	Pro	Phe	Thr	Cys	His 320
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30	Phe	Arg	Ala	Tyr 340	Leu	Ile	Gly	Gly	Leu 345	Met	Ala	Phe	Leu	Leu 350	Leu	Ala
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Ala Ile Gln Trp Asp Gly Asp Phe Thr Glu Gln Ala Gln Cys Ala Lys

DOYNERS DEEDED

515 520 525

Thr Lys Phe Trp Lys Lys Val Arg Tyr His Met Pro Pro Arg Arg Tyr 530

5 Pro Ala Ser Pro Pro Val Gln Leu Leu Gly His Thr Pro Arg Ile Pro 545 550

Gly

10